IN THE CLAIMS

Please amend the claims as follows:

Claim 1 (Currently Amended): A purified protein, comprising characterized in that:

- a) it has at least 40% identity, over its entire sequence, with the Pks13 protein of M. tuberculosis (SEQ ID NO: 1); and
- b) it has an acyltransferase domain (pfam00698), a keto acyl synthase domain (pfam02801 or pfam00109), at least one acyl carrier protein domain (COG0331 or COG0304), and a thioesterase domain (COG3319 or pfam00975); wherein
- c) it catalyzes the purified protein catalyzes a Claisen condensation or malonic condensation between an acyl-CoA or acyl-AMP molecule and an acylmalonyl-CoA molecule.

Claim 2 (Currently Amended): The <u>purified</u> protein as elaimed in <u>of</u> claim 1, <u>wherein</u> the <u>purified</u> protein eharacterized in that it catalyzes a Claisen condensation or malonic condensation between:

a) an acyl-CoA molecule of formula I, or an acyl-AMP molecule of formula Ia:

$$R_1$$
 CH_2 COA (I) R_1 CH_2 CH_2

in which wherein R₁ is a chain comprising from 6 to 68 carbon atoms, which may contain comprise one or more C=C double bonds, and/or one or more cis/trans cis, trans, or

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<u>combination thereof</u>, <u>and/or</u> <u>and</u> which may carry one or more side groups chosen <u>selected</u> <u>from the group consisting of from -CH₃</u>, =O and -O-CH₃;

and

b) an acylmalonyl-CoA molecule of formula II:

in which wherein R_2 is a linear alkane comprising from 10 to 24 carbon atoms; so as to form a β -keto acyl intermediate of formula III, or a β -keto ester of formula IIIa:

$$R_1$$
 CH_2 C

in which wherein R_1 and R_2 are as defined above, and X_1 is an acceptor molecule.

Claim 3 (Currently Amended): The <u>purified</u> protein <u>of claim 1</u> as claimed in either one of claims 1 and 2, characterized in that it exhibits <u>comprising</u> at least 70% identity with the sequence SEQ ID No.: 1 from *Mycobacterium tuberculosis*.

Claim 4 (Currently Amended): The protein of claim 2, as claimed in either one of claims 1 and 2, characterized in that it exhibits further comprising at least 70% sequence identity with the sequence SEQ ID No.: 2 from Corynebacterium glutamicum.

Claim 5 (Currently Amended): An expression vector, characterized in that it comprises comprising a polynucleotide sequence encoding [[a]] the protein as claimed in any one of claims 1 to 4 of claim 1.

Claim 6 (Currently Amended): A host cell, characterized in that it is transformed with [[an]] the expression vector as claimed in of claim 5.

Claim 7 (Currently Amended): The host cell as claimed in of claim 6, characterized in that it wherein the host cell is a prokaryotic cell.

Claim 8 (Currently Amended): A method for obtaining a protein, wherein the protein comprises

- a) at least 40% identity, over its entire sequence, with the Pks13 protein of M. tuberculosis (SEQ ID NO: 1); and
- b) an acyltransferase domain (pfam00698), a keto acyl synthase domain (pfam02801 or pfam00109), at least one acyl carrier protein domain (COG0331 or COG0304), and a thioesterase domain (COG3319 or pfam00975); wherein
- c) the purified protein catalyzes a Claisen condensation or malonic condensation
 between an acyl-CoA or acyl-AMP molecule and an acylmalonyl-CoA molecule, comprising

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as claimed in any one of claims 1 to 4, characterized in that it comprises:

- [[-]] culturing [[a]] the host cell as claimed in either one of of claim 6 elaims 6 and 7; and
 - [[-]] purifying said the protein from said the culture.

Claim 9 (Currently Amended): Method A method for inhibiting the biosynthesis of the a mycolata envelope in a bacterium, characterized in that it comprises comprising inhibiting, in said the bacterium bacteria, the expression or the activity of [[a]] the protein as claimed in any one of claims 1 to 4 of claim 1, thereby inhibiting the mycolata envelope biosynthesis.

Claim 10 (Currently Amended): The use of a protein as claimed in any one of claims

1 to 4, for screening for antibiotics that are active on mycolata A method of screening for an
antibiotic against bacteria that must synthesize mycolic acids to be viable, comprising
obtaining a transformed bacterium capable of surviving without producing mycolic acids,
culturing the bacterium, on an medium comprising agar and a compound, to form colonies,
and observing the appearance of the colonies, such that if the morphology of the colonies
goes from a shiny smooth appearance to a rough appearance, the compound is an antibiotic.

Claim 11 (Currently Amended): The use as claimed in method of claim 10, for sereening for wherein the antibiotics that are active on bacteria that must synthesize mycolic acids to be viable are mycobacteria.

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Claim 12 (New): The purified protein of claim 1, wherein the purified protein catalyzes a Claisen condensation between the acyl-CoA molecule and the acylmalonyl-CoA molecule.

Claim 13 (New): The purified protein of claim 1, wherein the purified protein catalyzes a Claisen condensation between the acyl-AMP molecule and the acylmalonyl-CoA molecule.

Claim 14 (New): The purified protein of claim 1, wherein the purified protein catalyzes a malonic condensation between the acyl-CoA molecule and the acylmalonyl-CoA molecule.

Claim 15 (New): The purified protein of claim 1, wherein the purified protein catalyzes a malonic condensation between the acyl-AMP molecule and the acylmalonyl-CoA molecule.

Claim 16 (New): The purified protein of claim 2, wherein the purified protein catalyzes a Claisen condensation between the acyl-CoA molecule of formula I and the acylmalonyl-CoA molecule of formula II.

Claim 17 (New): The purified protein of claim 2, wherein the purified protein catalyzes a Claisen condensation between the acyl-AMP molecule of formula Ia and the acylmalonyl –CoA molecule of formula II.

Claim 18 (New): The purified protein of claim 2, wherein the purified protein catalyzes a malonic condensation between the acyl-CoA molecule of formula I and the acylmalonyl-CoA molecule of formula II.

Claim 19 (New): The purified protein of claim 2, wherein the purified protein catalyzes a malonic condensation between the acyl-AMP molecule of formula Ia and the acylmalonyl –CoA molecule of formula II.

Claim 20 (New): An expression vector comprising a polynucleotide sequence encoding the protein of claim 2.